

Variable	Mean	SD	Min	Max
Age	30.5	4.2	18	45
Gender	Male			
Marital status	Married			
Education	High school			
Occupation	Teacher			
Income	1000			
Health status	Good			
Smoking status	Non-smoker			
Alcohol consumption	None			
Exercise frequency	Weekly			
Stress level	Low			
Sleep quality	Good			
Dietary habits	Healthy			
Family size	2			
Religious beliefs	Religious			
Community involvement	Active			
Life satisfaction	High			
Work-life balance	Good			
Financial stability	Stable			
Healthcare access	Easy			
Environmental quality	Good			
Public services	Good			
Infrastructure	Good			
Transportation	Good			
Communication	Good			
Education system	Good			
Healthcare system	Good			
Justice system	Good			
Government services	Good			
Public safety	Good			
Environmental protection	Good			
Cultural heritage	Good			
Language	English			
Religion	Christianity			
Ethnicity	White			
Marital status	Married			
Education	High school			
Occupation	Teacher			
Income	1000			
Health status	Good			
Smoking status	Non-smoker			
Alcohol consumption	None			
Exercise frequency	Weekly			
Stress level	Low			
Sleep quality	Good			
Dietary habits	Healthy			
Family size	2			
Religious beliefs	Religious			
Community involvement	Active			
Life satisfaction	High			
Work-life balance	Good			
Financial stability	Stable			
Healthcare access	Easy			
Environmental quality	Good			
Public services	Good			
Infrastructure	Good			
Transportation	Good			
Communication	Good			
Education system	Good			
Healthcare system	Good			
Justice system	Good			
Government services	Good			
Public safety	Good			
Environmental protection	Good			
Cultural heritage	Good			
Language	English			
Religion	Christianity			
Ethnicity	White			

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	GGAAAGUCGG AACAUCCCC AUUUUUCCU GAAGAGACGA AGUGAUCAAG AGAUCCUUCU	60
	CUCACAAGUC GAAACUCAGA GUC	83
<210>	60	
<211>	73	
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<223>	nanocircle vector	
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	GAAAAGTGGG CTACAGGGAG GTACCAGGTA ATGTACCACA ACGTGTGTTT CTCTGGTCTG	60
	CTTCTCAGGA ATC	73
<210>	61	
<211>	73	
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<213> Artificial Sequence

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<223> circular monomer

<400> 61

gauuccugag aagcagacca gagaaacaca cguuguggua cauuaccugg uaccuccug 60

uaguccaguu uuc 73

<210> 62

<211> 41

<212> DNA

<213> Artificial Sequence

<220>

<223> oligomer

<400> 62

gagatgttcc gactttccga ctctgagttt cgacttgtga g 41

<210> 63

<211> 42

<212> DNA

<213> Artificial Sequence

<220>

<223> oligomer

<400> 63

agaaggatct cttgatcact tcgtctcttc agggaaagat gg 42

<210> 64

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> splint oligonucleotide

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aagtcggaac atctcccatc tttccctgaa 30

<210> 65

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> splint oligonucleotide

<400> 65
tcaagagatc cttctctcac aagtcgaaac 30

<210> 66

<211> 83

<212> RNA

<213> Artificial Sequence

<220>

<223> AS83 vector monomer

<400> 66
ggaaagucgg aacaucuccc aucuuucccu gaagagacga agugaucaag agauccuucu 60

cucacaaguc gaaacucaga guc 83

<210> 67

<211> 83

<212> RNA

<213> Artificial Sequence

<220>

<223> H83 vector monomer

<400> 67
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cucacaaguc gaaaccaacg guc 83

<210> 68

<211> 16
<212> RNA
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<220>

<223> nucleotides 1752-1767 from HIV-1 gag

<400> 68
uuguuggucc aaaaug

16

<210> 69

<211> 15

<212> RNA

<213> Artificial Sequence

<220>

<223> target for the AS83 ribozyme which includes sequences 64-73 of av
ocado sunblotch viroid RNA

<400> 69
ucugagucgg aaagg

15

<210> 70

<211> 38

<212> DNA

<213> Artificial Sequence

<220>

<223> half-length oligonucleotide

<400> 70
cgaaaactgg actacagggg ggtaccaggt aatgtacc

38

<210> 71

<211> 35

<212> DNA

<213> Artificial Sequence

<220>

<223> half-length oligonucleotide

<400> 71

acaacgtgtg tttctctggt ctgcttctca ggaat

35

<210> 72

<211> 14

<212> RNA

<213> Artificial Sequence

$\langle 220 \rangle$

<223> sequences of HIV-1 pol gene

<400> 72

cuguagucca ggaa

14

[illegible]

<210> 73

<211> 56

<212> DNA

<213> Artificial sequence

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<223> circular ssDNA library constituent oligonucleotide

$\langle 220 \rangle$

<221> misc_feature

<222> (9) .. (48)

<223> a, g, c, or t

<400> 73

ttcgtctgnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnntc tttag

56

<210> 74

<211> 47

<212> DNA

<213> Artificial Sequence

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<220>

<223> splint

<400> 77

aactcctcag tcaatgtggt gcca

24

<210> 78

<211> 58

<212> DNA

<213> Artificial Sequence

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58

<210> 79

<211> 45

<212> DNA

<213> Artificial Sequence

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<223> circular ssDNA

<400> 79

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<210> 80

<211> 58

<212> DNA

<213> Artificial Sequence

<220>

<223> circular ssDNA

<400> 80

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58

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gactgaggag ttcgtctgtc tttcagtttc gtcct

35

<210> 85

<211> 28

<212> DNA

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cacggactca tcagaatggc aacacatt

28

<210> 86

<211> 58

<212> DNA

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<400> 86

gacactggag ttcgtctggt aaagtatggt gctacgactt ctttatttac cacgatgc

58

<210> 87

<211> 45

<212> DNA

<213> Artificial Sequence

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<400> 87

agaaagtgtt tcgtcctcac ggactcatca gagagcgttc actct

45

<210> 88

<211> 0
<212> DNA
<213> Skipped Sequence

<400> 88
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<210> 89
<211> 44
<212> DNA
<213> Artificial Sequence

<220>

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<400> 89
agaaagtgtg tcgtcctcac ggactcatca gagagcgttc actc

44

<210> 90

<211> 60

<212> DNA

<213> Artificial Sequence

<220>

<223> circular ssDNA

<220>

<221> misc_feature

<222> (11)..(50)

<223> a, g, c, or t

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agttcgtctg nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn agaaagtgtt

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<210> 91

<211> 43

<212> DNA

<213> Artificial Sequence

<220>

<223> circular ssDNA

<400> 91

tcgtcctcac ggactcatca gagagcggtc actctgacac tgg

43

<210> 92

<211> 35

<212> DNA

<213> Artificial Sequence

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<223> circular ssDNA

<400> 92

gacactggag ttcgtctgag aaagtgtttc gtcct

35

<210> 93

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> circular ssDNA

<400> 93

cacggactca tcagagagcg ttcactct

28

<210> 94

<211> 29

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 94

aggtcgacta tggagaaaaa aatcactgg

29

<210>	95	
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<210>	96	
<211>	81	
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	aatggagaaa aaaatcactg g	81
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<220>		

<221> misc_feature
<222> (19)..(58)
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<400> 101
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<210> 102
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<212> DNA
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<210> 103
<211> 0

<212> DNA
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000
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<210> 104
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<223> E1 motif I

<400> 104
gcaacgaatc agactctttc ggtgacattg cccagtttat 40

<210> 105
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<212> DNA

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<223> E38 motif II

<400> 105

gycacgatct gaatagtcgt tcatcctyag cggtagcgaa

40

<210> 106

<211> 0

<212> DNA

<213> Skipped Sequence

<400> 106

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<210> 107

<211> 0

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<400> 107

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<210> 108

<211> 40

<212> DNA

<213> Artificial Sequence

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<223> nucleotide sequence containing E15 motif III

<400> 108

gcttaaggat taattgcatg ttattcttta ggagcctcga

40

<210> 109

<211> 40

<212> DNA
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<220>

<223> nucleotide sequence containing E15 motif III

<400> 109
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<210> 110

<211> 0

<212> DNA

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<210> 111

<211> 40

<212> DNA

<213> Artificial Sequence

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<400> 111
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<210> 112

<211> 40

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<400> 112
tatgacgata ggattagacg tgtgggggta ttttcactac

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<210> 113
<211> 40
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<400> 113
gtctcactcg gaggaggagt ctgacaagat gggatgctgc 40

<210> 114
<211> 40
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<400> 114
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<210> 115
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gagtggtaag tactgggagc cactcacgac aacgaacaa 39

<210> 116
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<400> 116

gtgctattcg tggctatact gttaatgtgt cgcaccattc

40

<210> 117

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ggccccgttt aggtacaatc acatgtacta gcgttggtgt

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<210> 118

<211> 40

<212> DNA

<213> Artificial Sequence

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<223> selected nucleotide sequence

<400> 118

gtaggggttaa tatccttctc gtatgaccgt ggaagacgtc

40

<210> 119

<211> 40

<212> DNA

<213> Artificial Sequence

<220>

<223> selected nucleotide sequence

<400> 119

ggatgcgtag cgtaaagcgt tcgtatctcg aggtaagctt

40

<210> 120

<211> 40

<212> DNA

<213> Artificial Sequence

<220>

<223> selected nucleotide sequence

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<211> 103

<212> DNA

<213> Artificial Sequence

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tttcagtttc gtcctcacgg actcatcaga atggcaacac att 103

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<211> 103

<212> DNA

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<223> E15

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tttcagtttc gtcctcacgg actcatcaga atggcaacac att 103

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<211> 103

<212> DNA

<213> Artificial Sequence

<220>
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<220>
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 aaagugucaga gcg 13
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 <211> 103
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<220>
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 <400> 125
 agagugaacg cucucugaug aguccgugag gacgaaacac uuucugcauc gugguaaaaua 60
 aagaagucgu agcaacauac uuuaccagac gaacuccagu guc 103
 <210> 126
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 <212> RNA
 <213> Artificial Sequence

<220>
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<400> 126
agagugaacg cucucugaug aguccgugag gacgacacac uuucugcauc gugguaaaaua 60
aagaagucgu agcaacauac uuuaccagac gaacuccagu guc 103

<210> 127

<211> 63

<212> RNA

<213> Artificial Sequence

<220>

<223> short marA

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guc 63

<210> 128

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> probe

<400> 128
ctccctgtag tccagttttc g 21

<210> 129

<211> 73

<212> DNA

<213> Artificial Sequence

<220>

<223> DNA template encoding self-processing haripin ribozyme

<400> 129
ttcctgagaa gtcaaccaga gaaacacacg ttgtggtaca ttacctgcta cctccctgta 60
gtccagtttt cga 73